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INTRODUCTION

Two genes for hereditary breast cancer, BRCA1 and BRCA2, have now been identified (Miki et al, 1994, Wooster et al, 1995). The lifetime risk for breast cancer exceeds 80% in carriers of mutations in either of these genes (Ford et al, 1994). It is now possible to identify families with mutations in these two genes and genetic testing is underway in the laboratories of the principal investigator and the co-investigator.

The risk of second primary breast cancer in women with BRCA1 or BRCA2 mutations is high. Up to 60% of carriers will develop a contralateral cancer if they survive the initial cancer (Ford et al, 1994). Because of this very high risk, some women opt for prophylactic mastectomy. Other women are requesting either unilateral or bilateral mastectomy at the time of diagnosis in the hope that this will diminish their risk for subsequent cancers. There is no information yet on how these different treatments impact upon patient survival. Our objective is to establish whether women with hereditary breast cancer benefit from more extensive surgery than simple lumpectomy.

Tamoxifen use has been associated with a 39% reduction in contralateral breast cancer (Early Breast Cancer Trialists Collaborative Group). Because of the high risk of contralateral breast cancer in BRCA1 and BRCA2 carriers, it is of interest to establish if this drug is useful in reducing second primary breast cancers in this subgroup. Furthermore, if the rates of contralateral breast cancer are reduced substantially, there will be compelling reasons to believe that tamoxifen has potential as a chemopreventive agent in healthy gene carriers.

Similarly, ovarian ablation was associated with a 26% reduction in mortality from breast cancer in women under 50 with stage 1 or stage 2 disease (Early Breast Cancer Trialists Collaborative Group). From our early experience, 40% of women with breast cancer and BRCA1 mutations undergo oophorectomy at or near the time of mastectomy, primarily as a preventive measure against ovarian cancer. This provides us with the opportunity to ask whether or oophorectomy influences overall mortality from breast cancer and whether it reduces the incidence of contralateral tumours.

Our objective is to establish whether women with hereditary breast cancer benefit from more extensive surgery than simple lumpectomy.

BODY

Hypothesis:

Our main hypothesis is that mastectomy will be associated with a greater survival than lumpectomy in women diagnosed with stage I or stage II breast cancer, and who carry a BRCA1 or BRCA2 mutation. Our subhypotheses are: 1) treatment with tamoxifen reduces the risk of death from cancer and of contralateral breast cancer in this subgroup, and 2) that oophorectomy reduces the risk of death from cancer and of contralateral breast cancer in this subgroup.

Technical Objectives:

- 1) To identify 500 women, living or deceased, diagnosed with stage I or stage II breast cancer, from 1975 to the present, at age 65 or below, and who belong to a family documented to carry a BRCA1 or a BRCA2 mutation.
- 2) To review the medical record of these 500 women and to document the initial presentation and treatment of the breast cancer, including tumour size, stage, grade, the surgical procedure performed, the use of adjuvant chemotherapy, radiotherapy, tamoxifen, and oophorectomy.
- To obtain follow-up information on these women, including age and cause of death, age of local recurrence, if any, and age of contralateral breast cancer, if any.
- 4) To perform a survival analysis on this historical cohort using a proportional hazards

model, adjusting for size, stage and grade of tumor, number of nodes involved, the age of diagnosis and the estrogen and progesterone receptor status. The principal predictive variable of interest will be lumpectomy versus mastectomy (unilateral and bilateral). Additional variables of interest will be the use of tamoxifen (ever/never use and duration of use) and a history of bilateral oophorectomy (yes/no; date). We will calculate relative risks for survival based on the type of surgery, adjusting for the other predictive variables.

Statement of Work:

Task 1. Years 1-3:

We will to identify 500 women with breast cancer from a family for which a BRCA1 or a BRCA2 mutation has been identified during the three year period of this study. These women will be living or deceased, will have been diagnosed with stage I or II breast cancer in 1975 or thereafter, at the age of 65 or below.

Task 2. Years 1-3:

We will review the medical histories of these women and obtain information relating to surgical and medical treatment of the first breast cancer. We agree to obtain follow-up information on these women regarding the date and cause of death, breast cancer recurrence in the same and opposite breast.

Task 3. Years 2-3:

We will perform a statistical analysis on this data set in order to estimate the protective effect of mastectomy vs oophorectomy in this cohort, and evaluate whether there is any protection associated with tamoxifen use or with oophorectomy.

Experimental Methods

A historical cohort approach will be employed. 500 women with hereditary breast cancer, diagnosed since 1975, will be identified be identified by review of the pedigrees of families with BRCA1 or BRCA2 mutations. The medical charts of all of the breast cancer cases in the family, living and dead, will be reviewed. We will record the date of diagnosis, the type of surgical treatment performed (lumpectomy, unilateral mastectomy, or bilateral mastectomy), as well as the pathological stage and tumour grade, the use of adjuvant chemotherapy, radiotherapy, tamoxifen and oophorectomy.

We will compare the survival experiences of the women treated with the different surgical procedures, controlling for stage, nodal status, grade and the use of chemotherapy and radiotherapy. We will also evaluate additional survival benefits associated with oophorectomy and tamoxifen use in this cohort. The three principal endpoints will be local recurrence, contralateral breast cancer and death.

Progress To Date

As of October 2000, we have identified all eligible breast cancer cases within eligible families from eight clinical centres. We have enrolled 442 study subjects in this study, including 291 women with BRCA1 mutations, 139 women with BRCA2 mutations, and 11 women with BRCA1 and BRCA2 mutations identified within the family. We are continuing to collect medical records and complete follow-up on the remaining eligible cases. Thus, study subject recruitment and data collection is 90% complete. We have completed the medical chart review of all of these patients and have been able to establish the year and place of diagnosis, the stage and grade of the tumour, the surgical treatment performed, and the use of ovarian ablation, chemotherapy or oophorectomy. We have not performed a complete survival analysis on this cohort yet because data collection is ongoing. The details of the 442 patients are presented in Tables 1 to 3.

Table 1. Description of Study Subjects

Total number of cases in study	442
Centre:	
University of Toronto	94
Creighton	131
Pennsylvania	51
British Columbia	19
Chicago	65
Montreal	33
McGill	19
Beth Israel Medical Center	30
BRCA1	291
BRCA2	139
BRCA1/2	11
Mean age of diagnosis	42.0 years
Mean duration of follow-up	8.0 years

Table 2. Stage Distribution of 442 Tumours

		Number	Deceased (%)	mean age of dx
Stage	I	201	28 (13.9)	42.5
	II	241	48 (19.9)	41.6
Size	< 2cm	280	43 (15.4)	42.2
	> 2cm	160	33(20.6)	41.7
	missing	2		
Age	<40	204	44 (21.6)	34.6
	>40	238	32 (13.4))	48.3
Nodal Status				
	negative	295	47 (15.9)	42.3
	positive	146	29 (19.9)	41.3
	missing	1		

Table 3. Treatment Distribution of 442 cases

	Number	Deceased (%)	mean age of dx
Surgery:			
lumpectomy	159	26 (16.4)	42.1
unilateral mastectomy	175	43 (24.6)	42.9
bilateral mastectomy	108	7 (6.5))	40.1
Radiotherapy			
yes no missing	188 247 7	31 (16.5) 38 (15.4)	41.9 50.0
Chemotherapy			
yes no missing	268 162 12	38 (14.2) 30 (18.5)	40.5 44.3
Tamoxifen			
yes no missing	129 281 32	11 (8.5) 50 (17.9)	45.5 40.6
Oophorectomy			
bilateral unilateral none missing	156 21 243 22	16 (10.3) 4 (19.0) 45(18.5)	43.2 44.7 41.1

Data Analysis

The data will be analyzed by the Cox proportional hazards model using the SAS package. The 500 cases will form a historical cohort. Entry into the cohort will defined at the time of diagnosis of first breast cancer. The data will be analyzed at the time the cohort of 500 women is assembled, and at periods of five and ten years thereafter. Patients will be followed until either death, loss to follow-up, or the date of data analysis.

A survival curve will be constructed for the overall cohort and for the following three subgroups: 1) women treated with mastectomy +/- radiotherapy; 2) women treated with unilateral mastectomy; 3) women treated with bilateral mastectomy. Patients may opt for further preventive surgeries following the initial treatment and may go from group 1) to group 2) or 3), or from group 2) to 3) during the time frame of this study. They will be treated in three ways: 1) they will remain in the initially assigned group; 2) they will be excluded from consideration; 3) they will be considered to change groups and will be analyzed as such using time-dependent covariates. The principal endpoint for the comparison of surgical groups will be death. The Cox regression analysis allows for the control of extraneous variables which are related to prognosis. To control for these prognostic factors, the following will be entered into the Cox model: tumour size (in cm); number of positive axillary nodes; age at diagnosis (years); menopausal status (pre, peri, post); ER status (+/-); PR status (+/-); use of radiotherapy (yes/no); use of adjuvant chemotherapy (yes/no); tamoxifen (yes/no, and duration of use in months); oophorectomy (yes/no). For those women who underwent oophorectomy more than one year after the treatment of the breast cancer, oophorectomy will be treated as a time-dependent covariate.

The analysis will be conducted separately for the BRCA1 and BRCA2 subgroups. It will

also be possible to estimate the mortality hazard ratio for BRCA1 carriers compared with BRCA2 carriers.

The use of tamoxifen will be studied as an independent prognostic factor. In this tamoxifen substudy there will be two endpoints: contralateral breast cancer incidence and death. Again, the data will be analyzed separately for the BRCA1 and BRCA2 carriers. The usefulness of tamoxifen will be assessed in the cohort as a whole and separately in the three subgroups listed above. Similarly a history of oophorectomy will be studied as an independent risk factor in the overall cohort and in the subgroups defined above.

The analysis will be completed in the first six months of 2001. It is the intent of the principal investigator to submit the final manuscript by July 2001. No additional funds are required to complete the data analysis.

KEY RESEARCH ACCOMPLISHMENTS

- Risk of contralateral breast cancer in hereditary breast cancer
- Risk factors for contralateral breast cancer
- Effect of breast cancer treatment on survival
- Description of pathological characteristics of large cohort of hereditary breast cancer tumours
- Survival estimates for hereditary breast cancer

REPORTABLE OUTCOMES

Presentations

Metcalfe, K., Brunet, J., Lynch, H., & Narod, S. "Non-synchronous Contralateral Breast Cancer in Breast Cancer Patients from *BRCA1* and *BRCA2* Families". Controversies in the Etiology, Detection and Treatment of Breast Cancer. University of Toronto. May, 1998.

Metcalfe, K., Lynch, H., Weber, B., Olivotto, I., Foulkes, W., Ghadirian, P., Olopade, F., & Narod, S. "Non-synchronous Contralateral Breast Cancer in Breast Cancer Patients from *BRCA1* and *BRCA2* families". Familial Cancer Seminar. University of Vermont. October, 1998.

Metcalfe, K., Lynch, H., Weber, B., Olivotto, I., Foulkes, W., Ghadirian, P., Olopade, F., & Narod, S. "Contralateral Breast Cancer in Patients from *BRCA1* and *BRCA2* Families". Annual Canadian Collaborative Group for Cancer Genetics Conference. Calgary Alberta. November, 1998.

Metcalfe, K. "Risk Factors for Contralateral Breast Cancer". Workshop on the Prevention of Breast and Ovarian Cancer in BRCA1 and BRCA2 Carriers. Interdepartmental Division of Oncology, Faculty of Medicine, University of Toronto, June, 2000.

Narod, S., Metcalfe, K., Lynch, H., Ghadirian, P., Foulkes, W, Olivotto, I., Tung, N., Olopade, O., Brunet, J., & Weber, B. "Surgical Treatment of Hereditary Breast Cancer". Era of Hope:

Department of Defense Breast Cancer Research Program Meeting, Atlanta, Georgia, June 2000.

Metcalfe, K. "Surgical Treatment of Hereditary Breast Cancer". Women's Health Research in Progress Rounds. The Centre for Research in Women's Health, Toronto, October, 2000.

Abstracts

Narod, S., Metcalfe, K., Lynch, H., Ghadirian, P., Foulkes, W, Olivotto, I., Tung, N., Olopade, O., Brunet, J., & Weber, B. "Surgical Treatment of Hereditary Breast Cancer". Era of Hope: Department of Defense Breast Cancer Research Program Meeting, Atlanta, Georgia, June 2000.

PERSONNEL RECEIVING SUPPORT

Kelly Metcalfe

Danny Vesprini

William Chiu

CONCLUSIONS

This preliminary data set confirms our ability to ascertain the relevant clinical data and outcomes on BRCA1/BRCA2 carriers. We are continuing to complete families to ensure that all eligible breast cancer cases are included in the data collection and analysis. We will complete our collection of data with these remaining cases. A crude univariate analysis of this data suggests that improved survival is associated with age of diagnosis over 40, small tumours, stage I tumours, bilateral mastectomy, oophorectomy, chemotherapy, and tamoxifen. However, this preliminary analysis has not been adjusted for covariates (age and other prognostic features) or for variable length of follow-up. These adjustments will be made on the final data set, using the methods described in Data Analysis, above. It will also be necessary to estimate the prognostic factors separately for the subgroups of women with BRCA1 mutations and BRCA2 mutations.

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Wooster R, Bignell G, Swift S, Lancaster J et al. Identification of the breast cancer susceptibility gene BRCA2. Nature 378, 789-92, 1995